9590 5709

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/037,591A

DATE: 07/10/2002 TIME: 13:05:38

Input Set : A:\37128C.txt

Output Set: N:\CRF3\07102002\J037591A.raw

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4 <110> APPLICANT: Medlock, Eugene
              Yeh, Richard
      5
              Silbiger, Scott M.
      6
      7
              Elliot, Gary S.
             Nguyen, Hung Q.
              Jing, Shuqian
    11 <120> TITLE OF INVENTION: IL-17 Like Molecules and Uses Thereof
    13 <130> FILE REFERENCE: 01017/37128C
    15 <140> CURRENT APPLICATION NUMBER: 10/037,591A
C--> 16 <141> CURRENT FILING DATE: 2002-06-24
    18 <150> PRIOR APPLICATION NUMBER: 09/886,404
    19 <151> PRIOR FILING DATE: 2001-06-21
    21 <150> PRIOR APPLICATION NUMBER: 09/810,384
    22 <151> PRIOR FILING DATE: 2001-03-16
    24 <150> PRIOR APPLICATION NUMBER: 60/266,159
    25 <151> PRIOR FILING DATE: 2001-02-02
    27 <150> PRIOR APPLICATION NUMBER: 60/213,125
    28 <151> PRIOR FILING DATE: 2000-06-22
    30 <160> NUMBER OF SEQ ID NOS: 24
    32 <170> SOFTWARE: PatentIn Ver. 2.0
    34 <210> SEQ ID NO: 1
    35 <211> LENGTH: 644
    36 <212> TYPE: DNA
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    40 <221> NAME/KEY: CDS
    41 <222> LOCATION: (159)..(641)
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    44 ctcaagtcac tccctaaaaa gacagtggaa ataaatttga ataaacaaaa caggcttgct 60
    46 gaaaataaaa tcaggactcc taacctgctc cagtcagcct gcttccacga ggcctgtcag 120
    48 tcagtgcccc acttgtgact gagtgtgcag tgcccagc atg tac cag gtg gtt gca 176
                                                  Met Tyr Gln Val Val Ala
    49
    50
    52 ttc ttg gca atg gtc atg gga acc cac acc tac agc cac tgg ccc agc
                                                                           224
    53 Phe Leu Ala Met Val Met Gly Thr His Thr Tyr Ser His Trp Pro Ser
                     10
    56 tgc tgc ccc agc aaa ggg cag gac acc tct gag gag ctg ctg agg tgg
                                                                           272
    57 Cys Cys Pro Ser Lys Gly Gln Asp Thr Ser Glu Glu Leu Leu Arg Trp
                                     30
    60 agc act gtg cct gtg cct ccc cta gag cct gct agg ccc aac cgc cac
                                                                           320
    61 Ser Thr Val Pro Val Pro Pro Leu Glu Pro Ala Arg Pro Asn Arg His
                                                      50
                                 45
    62
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64 cca gag tcc tgt agg gcc agt gaa gat gga ccc ctc aac agc agg gcc

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65 Pro Glu Ser Cys Arg Ala Ser Glu Asp Gly Pro Leu Asn Ser Arg Ala
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68 atc tcc ccc tgg aga tat gag ttg gac aga gac ttg aac cgg ctc ccc
                                                                      416
69 Ile Ser Pro Trp Arg Tyr Glu Leu Asp Arg Asp Leu Asn Arg Leu Pro
                   75
                                        80
72 cag gac ctg tac cac gcc cgt tgc ctg tgc ccg cac tgc gtc agc cta
                                                                      464
73 Gln Asp Leu Tyr His Ala Arg Cys Leu Cys Pro His Cys Val Ser Leu
              . 90
                                    95
76 cag aca ggc tcc cac atg gac ccc cgg ggc aac tcg gag ctg ctc tac
                                                                      512
77 Gln Thr Gly Ser His Met Asp Pro Arg Gly Asn Ser Glu Leu Leu Tyr
                               110
80 cac aac cag act gtc ttc tac cgg cgg cca tgc cat ggc gag aag ggc
                                                                      560
81 His Asn Gln Thr Val Phe Tyr Arg Arg Pro Cys His Gly Glu Lys Gly
       120
                           125
                                                                      608
84 acc cac aag ggc tac tgc ctg gag cgc agg ctg tac cgt gtt tcc tta
85 Thr His Lys Gly Tyr Cys Leu Glu Arg Arg Leu Tyr Arg Val Ser Leu
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88 gct tgt gtg tgt gtg cgg ccc cgt gtg atg ggc tag
89 Ala Cys Val Cys Val Arg Pro Arg Val Met Gly
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94 <211> LENGTH: 161
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96 <213> ORGANISM: Homo sapiens
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102 Tyr Ser His Trp Pro Ser Cys Cys Pro Ser Lys Gly Gln Asp Thr Ser
                                     25
105 Glu Glu Leu Leu Arg Trp Ser Thr Val Pro Val Pro Pro Leu Glu Pro
                                 40
            35
108 Ala Arg Pro Asn Arg His Pro Glu Ser Cys Arg Ala Ser Glu Asp Gly
                             55
        50
111 Pro Leu Asn Ser Arg Ala Ile Ser Pro Trp Arg Tyr Glu Leu Asp Arg
                                             75
                         70
114 Asp Leu Asn Arg Leu Pro Gln Asp Leu Tyr His Ala Arg Cys Leu Cys
                                         90
117 Pro His Cys Val Ser Leu Gln Thr Gly Ser His Met Asp Pro Arg Gly
                                    105
                100
120 Asn Ser Glu Leu Leu Tyr His Asn Gln Thr Val Phe Tyr Arg Arg Pro
           115
                                120
123 Cys His Gly Glu Lys Gly Thr His Lys Gly Tyr Cys Leu Glu Arg Arg
                            135
                                                140
127 Leu Tyr Arg Val Ser Leu Ala Cys Val Cys Val Arg Pro Arg Val Met
128 145
                        150
130 Gly
133 <210> SEQ ID NO: 3
134 <211> LENGTH: 1013
135 <212> TYPE: DNA
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Input Set : A:\37128C.txt
Output Set: N:\CRF3\07102002\J037591A.raw

	<213> ORGANISM: <220> FEATURE:	Mus musculus	
	<221> NAME/KEY:	CDS	
	<222> LOCATION:		
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		Val Ala Phe Leu Ala Met Ile Val Gly Thr His Thr	
145	-	5 10 15	
		atc cag gag ggc tgc agt cac ttg ccc agc tgc tgc 90	6
		Ile Gln Glu Gly Cys Ser His Leu Pro Ser Cys Cys	
149	20		
151	ccc agc aaa gag	caa gaa ccc ccg gag gag tgg ctg aag tgg agc tct 14	44
152	Pro Ser Lys Glu	Gln Glu Pro Pro Glu Glu Trp Leu Lys Trp Ser Ser	
153	35	40 45	
155	gca tct gtg tcc	ccc cca gag cct ctg agc cac acc cac cac gca gaa 19	92
156	Ala Ser Val Ser	Pro Pro Glu Pro Leu Ser His Thr His His Ala Glu	
157		55 60	
			40
160	Ser Cys Arg Ala	Ser Lys Asp Gly Pro Leu Asn Ser Arg Ala Ile Ser	
161		70 75 80	
			88
	Pro Trp Ser Tyr	Glu Leu Asp Arg Asp Leu Asn Arg Val Pro Gln Asp	
165		85 90 95	
			36
		Arg Cys Leu Cys Pro His Cys Val Ser Leu Gln Thr	
169	100		0.4
		5	84
	-	Asp Pro Leu Gly Asn Ser Val Pro Leu Tyr His Asn 120 125	
173	115		32
		Tyr Arg Arg Pro Cys His Gly Glu Gly Thr His	32
177	130	135 140	
			80
		Leu Glu Arg Arg Leu Tyr Arg Val Ser Leu Ala Cys	•
	145	150 155 160	
			27
		Pro Arg Val Met Ala	
185	_	165	
		cggttg ggagagaggg ccaggtgtac aatcaccttg ccaatgcggg 58	87
		tccaaa gccctacctg aagcagcagg ctcccgggac aagatggagg 64	
191	acttggggag aaac	totgac ttttgcactt tttggaagca cttttgggaa ggagcaggtt 70	07
		agagga tgctgttgtg gcatttctac tcaggaacgg actccaaagg 76	
195	cctgctgacc ctgg	aageca tacteetgge teettteeee tgaateeeee aacteetgge 82	27
197	acaggcactt tctc	cacctc tececettig cettitgtig tgtttgttig tgcatgccaa 88	87
		aggtgt aattgccttg aaggatggtt ctgaggtgaa agctgttatc 94	
		ttatcc aaataaacat ctgtgtttaa aaaaaaaaaa aaaaaaaaaa	
203	aaaaaa	10	013
206	<210> SEQ ID NO	: 4	
207	<211> LENGTH: 1	69	

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Input Set : A:\37128C.txt

Output Set: N:\CRF3\07102002\J037591A.raw

208 <212> TYPE: PRT 209 <213> ORGANISM: Mus musculus 211 <400> SEQUENCE: 4 212 Met Tyr Gln Ala Val Ala Phe Leu Ala Met Ile Val Gly Thr His Thr 5 215 Val Ser Leu Arg Ile Gln Glu Gly Cys Ser His Leu Pro Ser Cys Cys 20 25 218 Pro Ser Lys Glu Gln Glu Pro Pro Glu Glu Trp Leu Lys Trp Ser Ser 35 40 221 Ala Ser Val Ser Pro Pro Glu Pro Leu Ser His Thr His His Ala Glu 55 224 Ser Cys Arg Ala Ser Lys Asp Gly Pro Leu Asn Ser Arg Ala Ile Ser 70 227 Pro Trp Ser Tyr Glu Leu Asp Arg Asp Leu Asn Arg Val Pro Gln Asp 90 230 Leu Tyr His Ala Arg Cys Leu Cys Pro His Cys Val Ser Leu Gln Thr 105 100 233 Gly Ser His Met Asp Pro Leu Gly Asn Ser Val Pro Leu Tyr His Asn 120 234 115 236 Gln Thr Val Phe Tyr Arg Arg Pro Cys His Gly Glu Glu Gly Thr His 135 237 130 239 Arg Arg Tyr Cys Leu Glu Arg Arg Leu Tyr Arg Val Ser Leu Ala Cys 150 242 Val Cys Val Arg Pro Arg Val Met Ala 165 246 <210> SEQ ID NO: 5 247 <211> LENGTH: 155 248 <212> TYPE: PRT 249 <213> ORGANISM: Homo sapiens 251 <400> SEQUENCE: 5 252 Met Thr Pro Gly Lys Thr Ser Leu Val Ser Leu Leu Leu Leu Ser 10 256 Leu Glu Ala Ile Val Lys Ala Gly Ile Thr Ile Pro Arg Asn Pro Gly 25 20 259 Cys Pro Asn Ser Glu Asp Lys Asn Phe Pro Arg Thr Val Met Val Asn 262 Leu Asn Ile His Asn Arg Asn Thr Asn Thr Asn Pro Lys Arg Ser Ser 55 265 Asp Tyr Tyr Asn Arg Ser Thr Ser Pro Trp Asn Leu His Arg Asn Glu 70 268 Asp Pro Glu Arg Tyr Pro Ser Val Ile Trp Glu Ala Lys Cys Arg His 85 90 271 Leu Gly Cys Ile Asn Ala Asp Gly Asn Val Asp Tyr His Met Asn Ser 100 105 274 Val Pro Ile Gln Glu Ile Leu Val Leu Arg Arg Glu Pro Pro His 115 120 277 Cys Pro Asn Ser Phe Arg Leu Glu Lys Ile Leu Val Ser Val Gly Cys 130 135 280 Thr Cys Val Thr Pro Ile Val His His Val Ala

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Output Set: N:\CRF3\07102002\J037591A.raw

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281 145
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285 <211> LENGTH: 117
286 <212> TYPE: PRT
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289 <400> SEQUENCE: 6
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293 Ala Gln Arg Lys Cys Glu Val Asn Leu Gln Leu Trp Met Ser Asn Lys
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                                     25
296 Arg Ser Leu Ser Pro Trp Gly Tyr Ser Ile Asn His Asp Pro Ser Arg
299 Ile Pro Val Asp Leu Pro Glu Ala Arg Cys Leu Cys Leu Gly Cys Val
         50
302 Asn Pro Phe Thr Met Gln Glu Asp Arg Ser Met Val Ser Val Pro Val
303 65
305 Phe Ser Gln Val Pro Val Arg Arg Leu Cys Pro Pro Pro Arg
308 Thr Gly Pro Cys Arg Gln Arg Ala Val Met Glu Thr Ile Val Ala Gly
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311 Cys Thr Cys Ile Phe
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315 <210> SEQ ID NO: 7
316 <211> LENGTH: 117
317 <212> TYPE: PRT
318 <213> ORGANISM: Homo sapiens
320 <400> SEQUENCE: 7
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325 Ala Gln Arg Lys Cys Glu Val Asn Leu Gln Leu Trp Met Ser Asn Lys
328 Arg Ser Leu Ser Pro Trp Gly Tyr Ser Ile Asn His Asp Pro Ser Arg
331 Ile Pro Val Asp Leu Pro Glu Ala Arg Cys Leu Cys Leu Gly Cys Val
                             55
334 Asn Pro Phe Thr Met Gln Glu Asp Arg Ser Met Val Ser Val Pro Val
                         70
                                             75
337 Phe Ser Gln Val Pro Val Arg Arg Leu Cys Pro Pro Pro Pro Arg
340 Thr Gly Pro Cys Arg Gln Arg Ala Val Met Glu Thr Ile Ala Val Gly
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343 Cys Thr Cys Ile Phe
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347 <210> SEQ ID NO: 8
348 <211> LENGTH: 197
349 <212> TYPE: PRT
350 <213> ORGANISM: Homo sapiens
352 <400> SEQUENCE: 8
353 Met Thr Leu Leu Pro Gly Leu Leu Phe Leu Thr Trp Leu His Thr Cys
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VERIFICATION SUMMARY

PATENT APPLICATION: US/10/037,591A

DATE: 07/10/2002 TIME: 13:05:39

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L:16 M:271 C: Current Filing Date differs, Replaced Current Filing Date